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(71) Applicant (for all designated States except US): GREENOVA-TION PFLANZENBIOTECHNOLOGIE GMBH [DE/DE]; Sonnenstrasse 5, D-79104 Freiburg im Breisgau (DE).

(72) Inventors; and

(75) Inventors/Applicants (for US only): BEYER. Peter [DE/DE]; In der Etzmatt 10, D-79423 Heitersheim (DE). POTRYKUS, Ingo [DE/CH]; Im Stigler 54, CH-4312 Magden (CH).

JOACHIM STÜRKEN PATENTANWALTSGE-(74) Agent: SELLSCHAFT MBH; Engesserstrasse 4b, D-79108 Freiburg im Breisgau (DE).

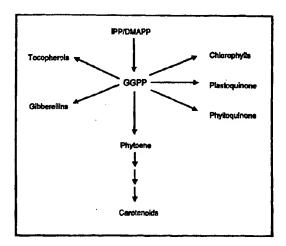
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(54) Title: METHOD FOR IMPROVING THE AGRONOMIC AND NUTRITIONAL VALUE OF PLANTS



(57) Abstract

The present invention provides means and methods of transforming plant cells, seeds, tissues or whole plants in order to yield transformants capable of expressing all enzymes of the carotenoid biosynthesis pathway that are essential for the targeted host plant to accumulate carotenes and/or xanthophylls of interest. The present invention also provides DNA molecules designed to be suitable for carrying out the method of the invention, and plasmids or vector systems comprising said molecules. Furthermore, the present invention provides transgenic plant cells, seeds, tissues and whole plants that display an improved nutritional quality and contain such DNA molecules and/or that have been generated by use of the methods of the present invention.

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Claims:

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- An isolated DNA molecule comprising a nucleotide sequence providing one or more expression cassettes capable of directing production of one or more enzymes specific for the carotenoid biosynthesis pathway selected from the group consisting of:
 - phytoene synthase derived from plants, fungi or bacteria,
 - phytoene desaturase derived from plants, fungi or bacteria,
 - ζ-carotene desaturase derived from plants, and
 - lycopene cyclase derived from plants, fungi or bacteria,
- under the proviso that an expression cassette capable of directing production of phytoene synthase alone is excluded.
- 2. The DNA molecule according to claim 1, wherein said expression cassette comprises one or more genes or cDNAs coding for plant, fungi or bacterial phytoene synthase, plant, fungi or bacterial phytoene desaturase, plant ζ-carotene desaturase, or plant, fungi or bacterial lycopene cyclase, each operably linked to a suitable constitutive, inducible or tissue-specific promoter allowing its expression in plant cells, seeds, tissues or whole plants, under the proviso that an expression cassette comprising a gene or cDNA coding for phytoene synthase alone is excluded.

- 3. The DNA molecule according to claim 1 or 2, further comprising at least one selectable marker gene or cDNA operably linked to a constitutive, inducible or tissue-specific promoter sequence allowing its expression in plant cells, seeds, tissues or whole plants.
- The DNA molecule according to any of claims 1 to 3, wherein the nucleotide sequence coding for phytoene synthase originates from plants, preferably expressed under the control of a tissue-specific promoter.
- 5. The DNA molecule according to any of claims 1 to 4, wherein the nucleotide sequence coding for phytocne desaturase originates from bacteria and is fused with a suitable plastid transit peptide encoding sequence, both of which preferably are expressed under the control of a tissue-specific or constitutive promoter.

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- 6. The DNA molecule according to any of claims 1 to 5, wherein the nucleotide sequence coding for lycopene cyclase originates from plants, preferably expressed under the control of a tissue-specific or constitutive promoter.
- The DNA molecule according to any of claims 2 to 6, wherein the selectable marker gene or cDNA is hygromycin phosphotransferase under the control of a constitutive promoter.
 - 8. The DNA molecule according to any of claims 1 to 7, wherein said nucleotide sequence comprises functional expression cassettes for both phytoene synthase and bacterial or fungi phytoene desaturase.

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- 9. The DNA molecule according to any of claims 1 to 7, wherein said nucleotide sequence comprises a functional expression cassette for lycopene cyclase.
- 15 10. The DNA molecule according to claim 5 or 8, wherein said plastid transit peptide sequence is derived from the pea Rubisco small subunit (tp).
 - A plasmid or vector system comprising one or more DNA molecules according to any of claims 1 to 10.
 - 12. A plasmid or vector system according to claim 11, which is derived from Agrobacterium tumefaciens.
- 13. A transgenic plant cell, seed, tissue or whole plant that contains a DNA molecule25 according to any of claims 1 to 10.
 - 14. A transgenic plant cell, seed, tissue or whole plant according to claim 13, selected from the group consisting of eukaryotic alga, embryophytes comprising *Bryophyta*, *Pteridophyta* and Spermatophyta such as *Gymnospermae* and *Angiospermae*, the latter including *Magnoliopsida*, *Rosopsida*, and *Liliopsida* ("monocots").

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- 15. A transgenic plant cell, seed, tissue or whole plant according to claim 14, selected from the group consisting of grain seeds, with rice, wheat, barley, oats, amaranth, flax, triticale, rye, and corn being preferred; oil seeds, with *Brassica* seeds, cotton seeds, soybean, safflower, sunflower, coconut, and palm being preferred; other edible seeds or seeds with edible parts selected from the group consisting of pumpkin, squash, sesame, poppy, grape, mung beans, peanut, peas, beans, radish, alfalfa, cocoa, coffee, hemp; tree nuts, with walnuts, almonds, pecans, and chick-peas being preferred; potatoes, carrots, sweet potatoes, tomato, pepper, cassava, willows, oaks, elm, maples, apples, bananas; and ornamental flowers, with lilies, orchids, sedges, roses, buttercups, petunias, phlox, violets, and sunflowers being preferred.
- 16. A method of transforming plant cells, seeds, tissues or whole plants in order to yield transformants capable of expressing all enzymes of the carotenoid biosynthesis pathway necessary to produce carotenes and xanthophylls of interest, comprising the transformation of said plant cells, seeds, tissues or whole plants with one or more DNA molecules according to any of claims 1 to 10, or with a plasmid or vector system according to claim 11 or 12.
- 17. A method according to claim 16, wherein said host plant cells, seeds or tissues selected for transformation normally are carotenoid-free.
 - 18. A method according to claim 16, wherein said host plant cells, seeds or tissues selected for transformation normally contain carotenoids in amounts desired to enlarge or modify.
- 25 19. A transformed whole plant regenerated from transformants yielded according to any of claims 16 to 18, or parts thereof, selected from the group consisting of eukaryotic alga, embryophytes comprising Bryophyta, Pteridophyta and Spermatophyta such as Gymnospermae and Angiospermae, the latter including Magnoliopsida, Rosopsida, and Liliopsida ("monocots").

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20. A transformed whole plant or part thereof according to claim 19, selected from the group consisting of grain seeds, with rice, wheat, barley, oats, amaranth, flax, triticale, rye, and corn being preferred; oil seeds, with Brassica seeds, cotton seeds, soybean, safflower, sunflower, coconut, and palm being preferred; other edible seeds or seeds with edible parts selected from the group consisting of pumpkin, squash, sesame, poppy, grape, mung beans, peanut, peas, beans, radish, alfalfa, cocoa, coffee, hemp; tree nuts, with walnuts, almonds, pecans, and chick-peas being preferred; potatoes, carrots, sweet potatoes, tomato, pepper, cassava, willows, oaks, elm, maples, apples, bananas; and ornamental flowers, with lilies, orchids, sedges, roses, buttercups, petunias, phlox, violets, and sunflowers being preferred.

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A. CLASSIFICATION OF SUBJECT MATTER IPC 7 C12N15/52 C12N15/53 C12N15/82 C12N5/10 A01H5/00 According to International Patent Classification (IPC) or to both national classification and IPC **B. FIELDS SEARCHED** Minimum documentation searched (classification system followed by classification symbols) IPC 7 C12N A01H Documentation searched other than minimum documentation to the extent that such documents are included in the fields searched Electronic data base consulted during the international search (name of data base and, where practical, search terms used) EPO-Internal, BIOSIS, CHEM ABS Data C. DOCUMENTS CONSIDERED TO BE RELEVANT Citation of document, with indication, where appropriate, of the relevant passages Relevant to claim No. EP 0 872 554 A (HOFFMANN LA ROCHE) 1.11 21 October 1998 (1998-10-21) the whole document X BARTLEY GLENN E ET AL: "Two Arabidopsis 1,11 thaliana carotene desaturases, phytoene desaturase and zeta-carotene desaturase, expressed in Escherichia coli, catalyze a poly-cis pathway to yield pro-lycopene." EUROPEAN JOURNAL OF BIOCHEMISTRY, vol. 259, no. 1-2, January 1999 (1999-01), pages 396-403, XP000925505 ISSN: 0014-2956 the whole document -/--X Further documents are listed in the continuation of box C. Patent family members are listed in annex. Special categories of cited documents : T later document published after the International filing date or priority date and not in conflict with the application but cited to understand the principle or theory underlying the "A" document defining the general state of the art which is not considered to be of particular relevance invention "E" earlier document but published on or after the International "X" document of particular relevance; the daimed invention cannot be considered novel or cannot be considered to filing date document which may throw doubts on priority claim(s) or which is cited to establish the publication date of another citation or other special reason (as specified) involve an inventive step when the document is taken alone "Y" document of particular relevance; the claimed invention cannot be considered to involve an inventive step when the document is combined with one or more other such docu-ments, such combination being obvious to a person skilled *O" document referring to an oral disclosure, use, exhibition or other means document published prior to the international filing date but later than the priority date claimed "&" document member of the same patent family Date of the actual completion of the international search Date of mailing of the international search report 24 July 2000 07/08/2000 Name and mailing address of the ISA Authorized officer European Peterit Office, P.B. 5818 Petentiaan 2 NL – 2280 HV Rijswijk Tel. (+31-70) 340-2040, Tx. 31 651 epo ni,

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Kania, T



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